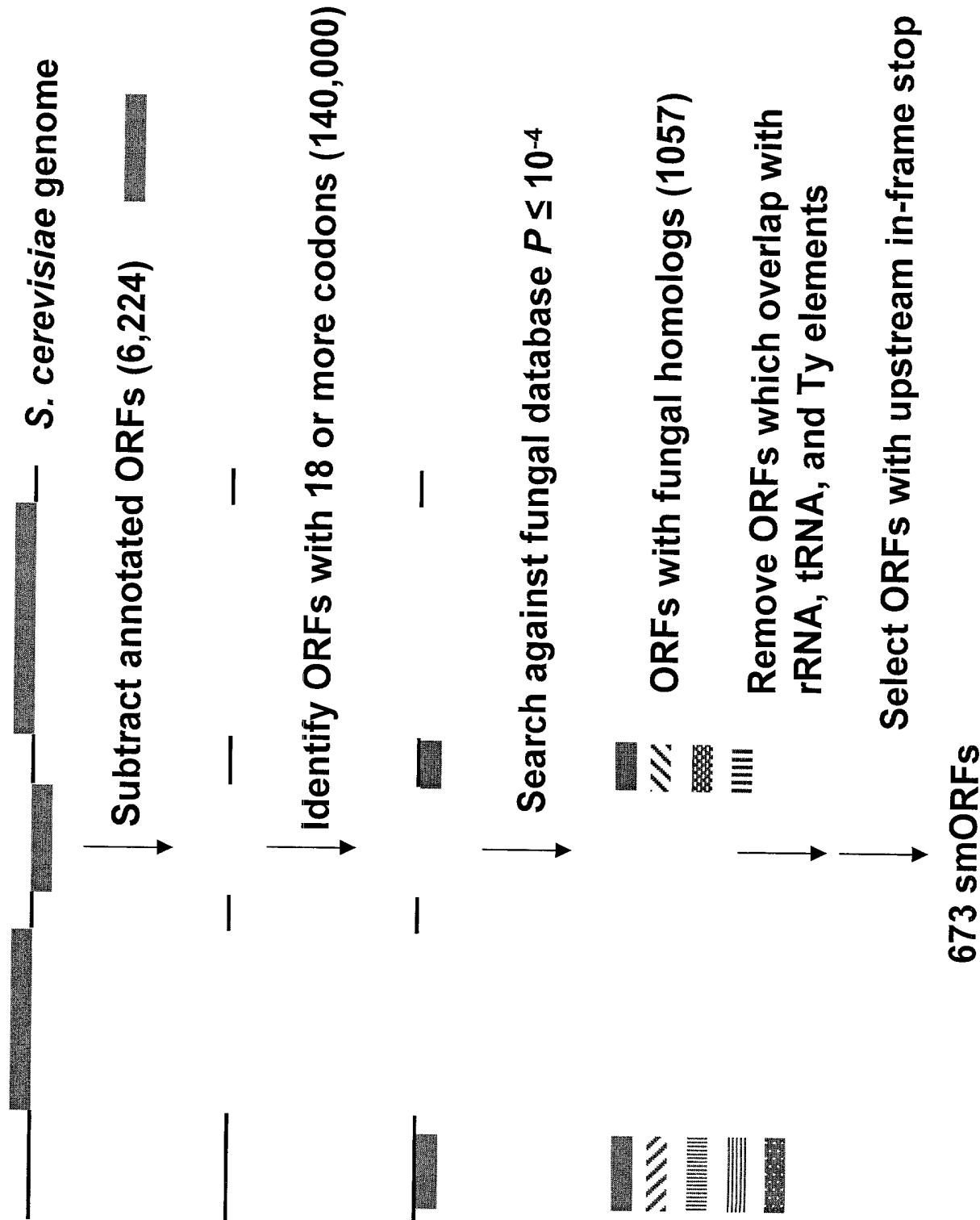


smORFs identification



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 INVENTOR(S): QIANDONG ZENG ET AL.
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Fig. 1

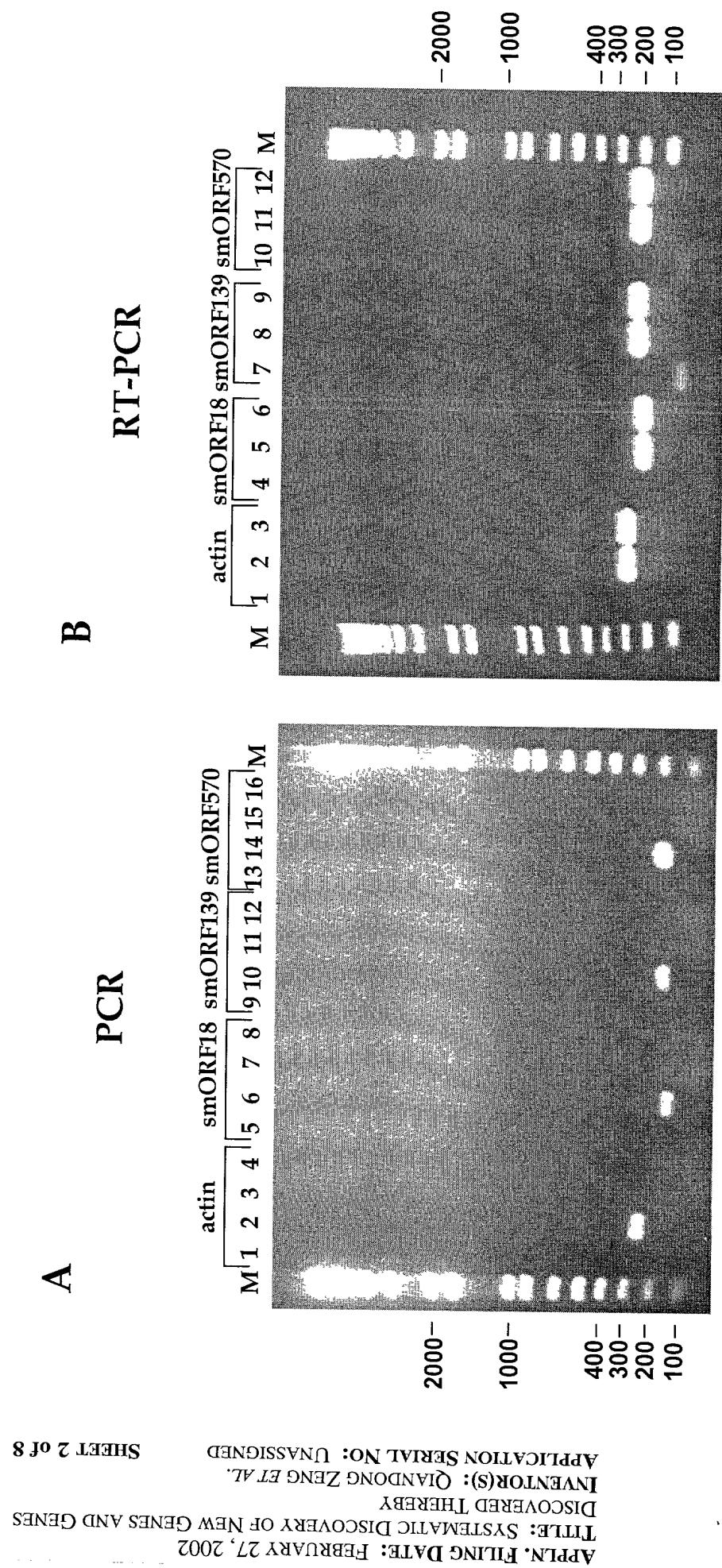
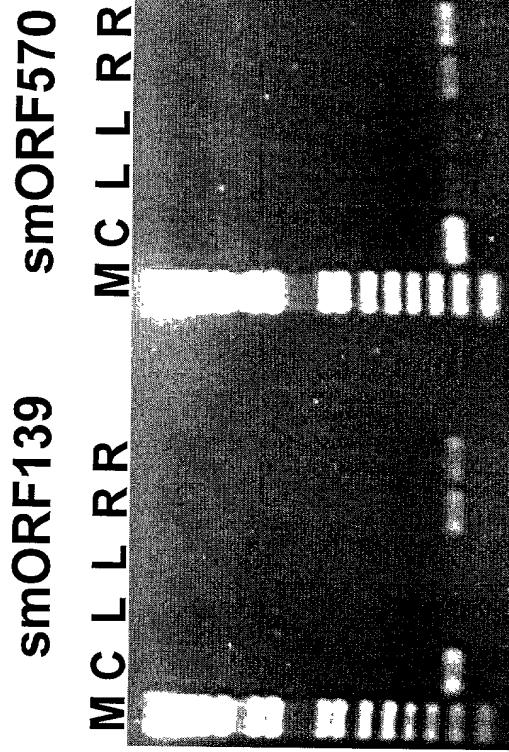
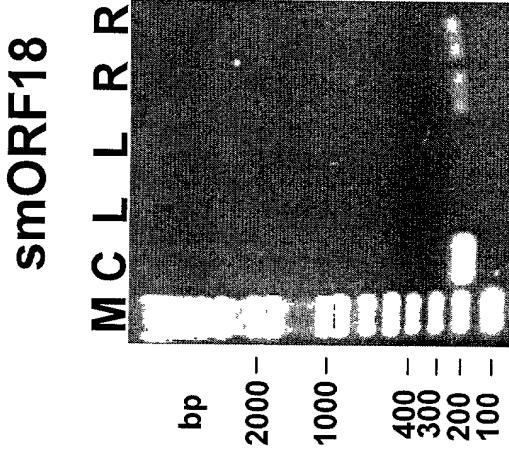
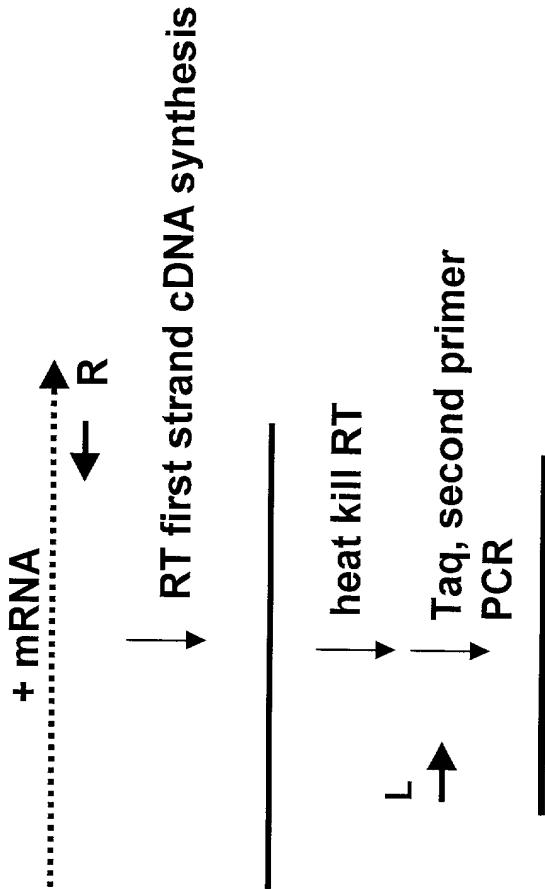


Fig. 2

Two step orientation-specific RT-PCR

C



1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18

Fig. 2

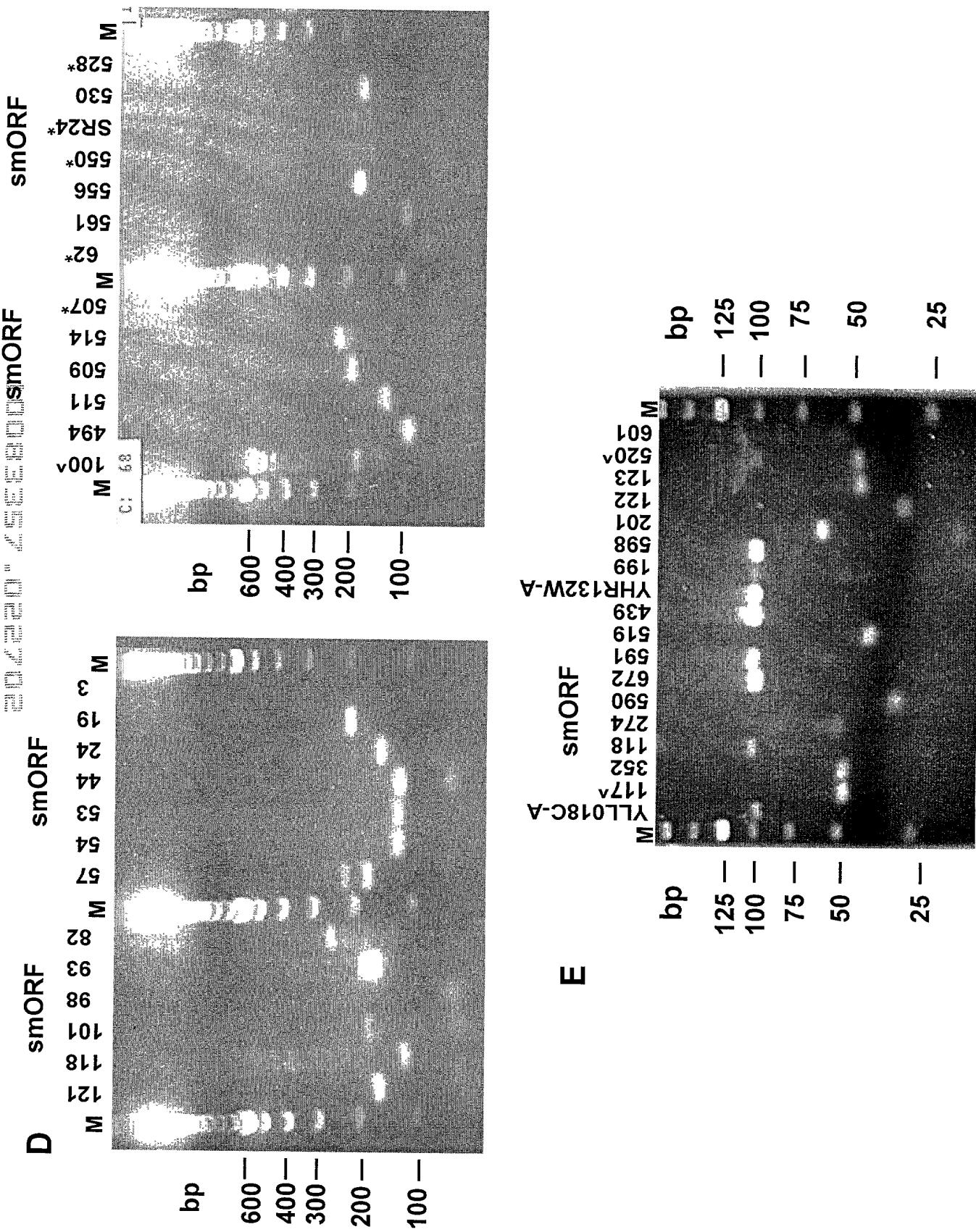


Fig. 2

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smORF18

Dm : MEE-----IHDVEVEIEDFYEYDIEEENYYYPCCPCGDFDIDKKEELIIGEEVATCPSSSLVIVYDPEMKAKEDIE--SALNEKLGDLKLEN
 Hs : MAY-----FHDEVEIEDFQYDIEDSHTYFYPCCPCGDNFSSIKEDIENGEDVATCPSSSLIIVKIVYDKDQEWCGETVPP--ABS----ANKELVIC
 Ce : MSV-----FHDEVEIEDFEEFDEEKDYIYYPCCPCGDPFVPIPREMILEMGEEDVAQCPSSCSLIRRIVYDDEDEVKLETIS--LSK----PIAEPV--
 Sp : MS-----FHDEVEIEDFADGTMYTTFPCPCGDRFEDIDOLGEDEEDVARDARCPSSCSLIRRIVYDDEDEVKLETIS--STA----PIIAAA
 Sc : MS-----T-YDEIEIEDMFEPEONYTYPCCPCGDRFPIYDDMFEGEKVAVCPSSSLMIDVVFEDKEDILAYYYEAGIHPE--PIAA--
 Ca : ME-----T-YDEIEIEDMFDPPVQCFQYPCCPCGDFRAIAYBMDNIGEDEIACPSSCSLMLVVFIFEPDILEYLEG
 Af : MADEAMISYDEIEIEDMFDPPNLQYYPCCPCGDRFEDIDRDGEDEIAVCPSSCSLMLVVFIFEVSDLPKDGNQPP--APCA--
 An : MADEAMISYDEIEIEDMFDANLQYYPCCPCGDFIAADLRYGEDEIAVCPSSCSLMLVVFQSDLPLKEEKKE--TEC--VSVA--VSYKA--

smORF139

Sc : MPKRLLSGLQKEVLTYRASIFTAHKPKENQVNFFVNYTHEFGKYRNLLRKFFLTIEELLRVRGKKKATFSLPHELLTNTI--
 Ca : MARLGSGLQRDVLIQYRKCIIRMVKKEIDHWRNEEEFAKYOHLPKKQENIEYLIRNGHRKLYNPKLQDI--
 Af : MARLGSGLQRDVLSYRKCLREIRKKPSETGRNFAYTRAEPKMSVSIKKFSAIEYLLRKGNRQIEVYSSSPGIEINTQ--
 Sp : MARLGSGLQRDVTFYRQLHAAKAKEQPYNTERWMAFVHQEFPKNQITISKRDFFYIETLLRVRGQRQYFAYSPFVWVDIIES

smORF570

Sc : -----SALENFRSLLQVLLLITCS-----SYHGGQWPSSLLIDYKNEHVAEWKMARVGERASPYVSLACLMAS----QFNS--
 Ca : KPSISPTNATKSALFNFSSLLQVSSLLITCTYMHSTAPLLDRNK-SGTSFWMKFARIGERLSPYVAKGCFIMAIEN----TSS--
 Bt : -----MSAIFNFSSLLIVLLLITCTYAYTSLAPSSLLDRNK-TGILGTFWKCARIGERKSPYVAVCCVMAFS----TLEQO--
 HS : -----MSAIFNFSSLLIVLLLITCTYAYTSLAPSSLLDRNK-TGILGTFWKCARIGERKSPYVAVCCVMAFS----TLEQO--
 Mm : -----MSAIFNFSSLLIVLLLITCTYAYTSLAPSSLLDRNK-TGILGTFWKCARIGERKSPYVAVCCVMAFS----TLEQO--
 Dm : -----NPSSALFNFHSLLIVLLLITCTYAYTSLAPSSLLDRNK-TGEMGTFWKLLARIGERKSPWVGAACTIMAFL----VLEFWS--
 Af : -----SQSALFNFSSLLIVFLLIICSTYAHSSMPGIMDRAQ--RGFETFWKCARIGERLSPYVVSLLCILMAVSTPRFVYVLLSNAY

1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40 41 42 43 44 45 46 47 48 49 50 51 52 53 54 55 56 57 58 59 60 61 62 63 64 65 66 67 68 69 70 71 72 73 74 75 76 77 78 79 80 81 82 83 84 85 86 87 88 89 90 91 92 93 94 95 96 97 98 99 100

smORF18-HA³

smORF18-HA³

wildtype

markers

213—
128—
85—

42.6—
31.2—

18—

8.5—

1 2 3 4

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Fig. 4

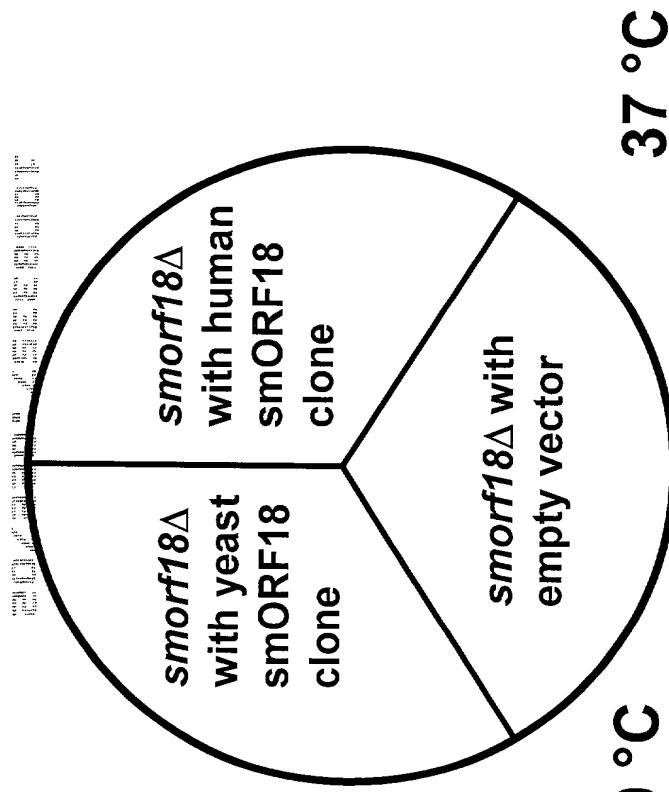


Fig. 5

smORF57 protein interaction map

